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| 1. Record Nr.           | UNISA996390488503316  |
| Autore                  | Ridpath George <d. 1726.>   |
| Titolo                  | Short-hand yet shorter: or The art of short-writing advanced in a more swift, easy, regular and natural method than hitherto [[electronic resource] ] : Whereby the former difficulties in placing the vowels are removed; they, the diphthongs and consonants, further contracted; the particles, pronouns, degrees of comparison, persons, moods, tenses, contrarieties, repetitions, sentences negative and interrogatory, are shortned. The rules are plain, easy to be remembred and applied to any short-hand, that such as have learned other authors may have hence a very considerable help to write more swiftly without altering their foundation. By George Ridpath |
| Pubbl/distr/stampa      | London, : printed for Andrew Bell, at the Cross-Keys in the Poultry near Stocks-Market, 1696  |
| Descrizione fisica      | [4], 43, [1] p., [2] leaves of plates (fold.) : tables  |
| Soggetti                | Shorthand   |
| Lingua di pubblicazione | Inglese   |
| Formato                 | Materiale a stampa  |
| Livello bibliografico   | Monografia  |
| Note generali           | Includes an errata at foot of p. 43.<br>Copy lacks A2.<br>Reproduction of the original in the University of London.   |
| Sommario/riassunto      | eebo-0169   |

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| 2. Record Nr.           | UNINA9910810663903321   |
| Autore                  | Amaratunga Dhammika <1956->   |
| Titolo                  | Exploration and analysis of DNA microarray and other high dimensional data / / Dhammika Amaratunga, Javier Cabrera, Ziv Shkedy, authors   |
| Pubbl/distr/stampa      | Hoboken, New Jersey : , : John Wiley & Sons, , 2014<br>©2014  |
| ISBN                    | 1-118-36450-3<br>1-118-36453-8  |
| Edizione                | [Second edition.]   |
| Descrizione fisica      | 1 online resource (346 p.)  |
| Collana                 | Wiley series in probability and statistics  |
| Altri autori (Persone)  | CabreraJavier<br>ShkedyZiv  |
| Disciplina              | 572/.636  |
| Soggetti                | Protein microarrays - Statistical methods   |
| Lingua di pubblicazione | Inglese   |
| Formato                 | Materiale a stampa  |
| Livello bibliografico   | Monografia  |
| Note generali           | Description based upon print version of record.   |
| Nota di bibliografia    | Includes bibliographical references and index.  |
| Nota di contenuto       | Cover; Title Page; Contents; Preface; Chapter 1 A Brief Introduction; 1.1 A Note on Exploratory Data Analysis; 1.2 Computing Considerations and Software; 1.3 A Brief Outline of the Book; 1.4 Data Sets and Case Studies; 1.4.1 The Golub Data; 1.4.2 The Mouse5 Data; 1.4.3 The Khan Data; 1.4.4 The Sialin Data; 1.4.5 The Behavioral Study Data; 1.4.6 The Spiked-In Data; 1.4.7 The APOAI Study; 1.4.8 The Breast Cancer Data; 1.4.9 Platinum Spike Data Set; 1.4.10 Human Epidermal Squamous Carcinoma Cell Line A431 Experiment; 1.4.11 Note: Public Repositories of Microarray Data; Chapter 2 Genomics Basics<br>2.1 Genes2.2 Deoxyribonucleic Acid; 2.3 Gene Expression; 2.4 Hybridization Assays and Other Laboratory Techniques; 2.5 The Human Genome; 2.6 Genome Variations and Their Consequences; 2.7 Genomics; 2.8 The Role of Genomics in Pharmaceutical Research and Clinical Practice; 2.9 Proteins; 2.10 Bioinformatics; Supplementary Reading; Chapter 3 Microarrays; 3.1 Types of Microarray Experiments; 3.1.1 Experiment Type 1: Tissue-Specific Gene Expression; 3.1.2 Experiment Type 2: Developmental Genetics; 3.1.3 Experiment Type 3: Genetic Diseases; 3.1.4 Experiment Type 4: Complex Diseases<br>3.1.5 Experiment Type 5: Pharmacological Agents3.1.6 Experiment Type 6: Plant Breeding; 3.1.7 Experiment Type 7: Environmental Monitoring; 3.2 A Very Simple Hypothetical Microarray Experiment; 3.3 |

A Typical Microarray Experiment; 3.3.1 Microarray Preparation; 3.3.2 Sample Preparation; 3.3.3 The Hybridization Step; 3.3.4 Scanning the Microarray; 3.3.5 Interpreting the Scanned Image; 3.4 Multichannel cDNA Microarrays; 3.5 Oligonucleotide Microarrays; 3.6 Bead-Based Arrays; 3.7 Confirmation of Microarray Results; Supplementary Reading and Electronic References

Chapter 4 Processing the Scanned Image

4.1 Converting the Scanned Image to the Spotted Image; 4.1.1 Gridding; 4.1.2 Segmentation; 4.1.3 Quantification; 4.2 Quality Assessment; 4.2.1 Visualizing the Spotted Image; 4.2.2 Numerical Evaluation of Array Quality; 4.2.3 Spatial Problems; 4.2.4 Spatial Randomness; 4.2.5 Quality Control of Arrays; 4.2.6 Assessment of Spot Quality; 4.3 Adjusting for Background; 4.3.1 Estimating the Background; 4.3.2 Adjusting for the Estimated Background; 4.4 Expression-Level Calculation for Two-Channel cDNA Microarrays

4.5 Expression-Level Calculation for Oligonucleotide Microarrays

4.5.1 The Average Difference; 4.5.2 A Weighted Average Difference; 4.5.3 Perfect Matches Only; 4.5.4 Background Adjustment Approach; 4.5.5 Model-Based Approach; 4.5.6 Absent-Present Calls; Supplementary Reading; Software Notes; Chapter 5 Preprocessing Microarray Data; 5.1 Logarithmic Transformation; 5.2 Variance Stabilizing Transformations; 5.3 Sources of Bias; 5.4 Normalization; 5.5 Intensity-Dependent Normalization; 5.5.1 Smooth Function Normalization; 5.5.2 Quantile Normalization; 5.5.3 Stagewise Normalization

5.5.4 Normalization of Two-Channel Arrays

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Sommario/riassunto

Praise for the First Edition "...extremely well written...a comprehensive and up-to-date overview of this important field." - Journal of Environmental Quality

Exploration and Analysis of DNA Microarray and Other High-Dimensional Data, Second Edition provides comprehensive coverage of recent advancements in microarray data analysis. A cutting-edge guide, the Second Edition demonstrates various methodologies for analyzing data in biomedical research and offers an overview of the modern techniques used in microarray technology to

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